

Fig. 1

1 AGCTCACAGCC 11  
 12 ATGGTTACCTTCAGCCACGTCTCCAGTCTGAGTCACTGGTTCCTCTTGCTGCTGCTGCTG 71  
 1 M V T F S H V S S L S H W F L L L L L L 20  
 72 AATCTGTTCTTGCCGGTAATATTTGCTATGCCTGAATCATACTCCTTCAACTGTCCCGAT 131  
 21 N L F L P V I F A M P E S Y S F N C P D 40  
 132 GGTGAATACCAGTCTAATGATGTCTGTTGCAAGACCTGTCCCTCAGGTACATTTGTCAAG 191  
 41 G E Y Q S N D V C C K T C P S G T F V K 60  
 192 GCGCCCTGCAAAATCCCCATACTCAAGGACAATGTGAGAAGTGTCAACCAGGAACATTC 251  
 61 A P C K I P H T Q G Q C E K C H P G T F 80  
 252 ACAGGGAAAGATAATGGCCTGCATGATTGTGAACTTTGCTCCACCTGTGATAAAGACCAG 311  
 81 T G K D N G L H D C E L C S T C D K D Q 100  
 312 AATATGGTGGCTGACTGTCTGCCACCAGTGACCGGAAATGCGAGTGCCAAATAGGTCTT 371  
 101 H M V A D C S A T S D R K C E C Q I G L 120  
 372 TACTACTATGACCCAAAATTTCCGGAATCATGCCGCCCATGTACCAAGTGTCCCAAGGA 431  
 121 Y Y Y D P K F P E S C R P C T K C P Q G 140  
 432 ATCCCTGTCTCCAGGAATGCAACTCCACAGCTAACACTGTGTGCAGTTTCTCTGTTTCA 491  
 141 I P V L Q E C N S T A N T V C S S S V S 160  
 492 AATCCCAGAAACTGGCTGTTCTCTACTGATGCTAATTGTCTTCTGTATCTGA 542  
 161 N P R N W L F L L M L I V F C I \* 177  
 543 AGAAGATAAAGGTTCTACAGATGGTGTCTGTAGCTTCCTTTTATTGCTGTGAAGAGAA 600  
 601 ACCATGGAGGCAACTCTTTCATTTTATTTTATTTTAAATGTCTTGAAGTTGATTGAAG 660  
 661 ACCAGGCTGGACTCAAACCTCACAGAGATCCGGACTAGGCACCTCTAATATAGGAAAAACAT 720  
 721 TGAATTGGGACTGGCTTACAGTTTCAGAAGTTCTGTCCATGATTATCATAGTGCGAAGCA 780  
 781 TGGAGGCACGGAGGCACACATGGTGTGAGAGAAGAGCTGAGAGTTCTGCATCTTGATCT 840  
 841 GCAAGCAATAAAAGGAGACTGTGTGCCACACTACACATAGCTTGAACATAGGAGACCTCA 900  
 901 AAGCCTGTCCCCACAGTGACAACTTCCTCCAACAAGGTATACCTCCTAATAATACCAT 960  
 961 TTCTTATGAGGCAAGCATTCAAACACATGAGTCTATGAGGGCCAAACCAATTCAAACCA 1020  
 1021 CACAGGTTAACAATTGCCCTCTGCAGCTCTCTGGTGGAGGCCCTCCTTGAGAGTAAGTAA 1080  
 1081 CAATTTAGATGAAGGCAAGTCTGGTATCAGGTCCAAAAGAACTCAGGATGAATGGTCC 1140  
 1141 ACTGTGGTTCTTATTAACATACTGAAGAACATGACCTCACCTTAGACTTCTCCACCTCAC 1200  
 1201 TGGCTTCCCTTCCCCTAGCTTCTCATTCCCAGGTAACCCTGCCATTTTTTGGTAATGTGC 1260  
 1261 CTTCTTGGTTCTTCTCTCTCTTCCCCCTCTCTTCTGGTCTTATTTCTCTTCTCTCCC 1320  
 1321 ACTCTCCACCAGCCGCTCTTAAGGCCTGAGTCAGTCTGACGGCCATGTTTAATCTACTA 1380  
 1381 CTTTCTCTCTGCTCTGGACTCATCCAGATGTCTCTGGCTGAGCTCTCCCTCTATCTACA 1440  
 1441 ATAAAACCTTCCCCCTAACAGAAATGGAACAGTTTTGCTCTCACTTTGTACATCTGGTG 1500  
 1501 CCTGAAACC 1509

Fig. 2

7F4 GPDGEY---QSNDVG CKTQPSGTFVKAPCK IPHTQQQCEKCHPGT FTGKDNGLHDGELCS 60  
mTNFR GPGGKYVHSKNNSIG GTCCHKGTYLVSQCP SPGRDTVCRECEKGT FTASQNYLRQCLSC 60

7F4 TQDKD--QNMVADGS ATSDRKCEC---QIG LYYDPKEPESGRPC TKCPQGIPVLQEGNS 120  
mTNFR TQKEMSQVEISPCQ ADKDTVCGCKENQFQ RYLSETHFQ--GVDC SPCFNGTVTIP-CKE 120

7F4 TANTVG 126  
mTNFR TQNTVG 126

7F4 TANTVG 126  
mTNFR TQNTVG 126



[REDACTED]



[REDACTED]



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1977-80	1978-80	1979-80	1980-81	1981-82	1982-83	1983-84	1984-85	1985-86	1986-87	1987-88	1988-89	1989-90	1990-91	1991-92	1992-93	1993-94	1994-95	1995-96	1996-97	1997-98	1998-99	1999-00	2000-01	2001-02	2002-03	2003-04	2004-05	2005-06	2006-07	2007-08	2008-09	2009-10	2010-11	2011-12	2012-13	2013-14	2014-15	2015-16	2016-17	2017-18	2018-19	2019-20	2020-21	2021-22	2022-23	2023-24	2024-25	2025-26	2026-27	2027-28	2028-29	2029-30	2030-31	2031-32	2032-33	2033-34	2034-35	2035-36	2036-37	2037-38	2038-39	2039-40	2040-41	2041-42	2042-43	2043-44	2044-45	2045-46	2046-47	2047-48	2048-49	2049-50	2050-51	2051-52	2052-53	2053-54	2054-55	2055-56	2056-57	2057-58	2058-59	2059-60	2060-61	2061-62	2062-63	2063-64	2064-65	2065-66	2066-67	2067-68	2068-69	2069-70	2070-71	2071-72	2072-73	2073-74	2074-75	2075-76	2076-77	2077-78	2078-79	2079-80	2080-81	2081-82	2082-83	2083-84	2084-85	2085-86	2086-87	2087-88	2088-89	2089-90	2090-91	2091-92	2092-93	2093-94	2094-95	2095-96	2096-97	2097-98	2098-99	2099-00	2100-01	2101-02	2102-03	2103-04	2104-05	2105-06	2106-07	2107-08	2108-09	2109-10	2110-11	2111-12	2112-13	2113-14	2114-15	2115-16	2116-17	2117-18	2118-19	2119-20	2120-21	2121-22	2122-23	2123-24	2124-25	2125-26	2126-27	2127-28	2128-29	2129-30	2130-31	2131-32	2132-33	2133-34	2134-35	2135-36	2136-37	2137-38	2138-39	2139-40	2140-41	2141-42	2142-43	2143-44	2144-45	2145-46	2146-47	2147-48	2148-49	2149-50	2150-51	2151-52	2152-53	2153-54	2154-55	2155-56	2156-57	2157-58	2158-59	2159-60	2160-61	2161-62	2162-63	2163-64	2164-65	2165-66	2166-67	2167-68	2168-69	2169-70	2170-71	2171-72	2172-73	2173-74	2174-75	2175-76	2176-77	2177-78	2178-79	2179-80	2180-81	2181-82	2182-83	2183-84	2184-85	2185-86	2186-87	2187-88	2188-89	2189-90	2190-91	2191-92	2192-93	2193-94	2194-95	2195-96	2196-97	2197-98	2198-99	2199-00	2200-01	2201-02	2202-03	2203-04	2204-05	2205-06	2206-07	2207-08	2208-09	2209-10	2210-11	2211-12	2212-13	2213-14	2214-15	2215-16	2216-17	2217-18	2218-19	2219-20	2220-21	2221-22	2222-23	2223-24	2224-25	2225-26	2226-27	2227-28	2228-29	2229-30	2230-31	2231-32	2232-33	2233-34	2234-35	2235-36	2236-37	2237-38	2238-39	2239-40	2240-41	2241-42	2242-43	2243-44	2244-45	2245-46	2246-47	2247-48	2248-49	2249-50	2250-51	2251-52	2252-53	2253-54	2254-55	2255-56	2256-57	2257-58	2258-59	2259-60	2260-61	2261-62	2262-63	2263-64	2264-65	2265-66	2266-67	2267-68	2268-69	2269-70	2270-71	2271-72	2272-73	2273-74	2274-75	2275-76	2276-77	2277-78	2278-79	2279-80	2280-81	2281-82	2282-83	2283-84	2284-85	2285-86	2286-87	2287-88	2288-89	2289-90	2290-91	229
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Fig. 4

peripheral blood leukocyte

colon

small intestine

ovary

testis

prostate gland

thymus

spleen

pancreas

kidney

skeletal muscle

liver

lung

placenta

brain

heart

Fig. 4  
peripheral blood leukocyte  
colon  
small intestine  
ovary  
testis  
prostate gland  
thymus  
spleen  
pancreas  
kidney  
skeletal muscle  
liver  
lung  
placenta  
brain  
heart

Fig. 5A

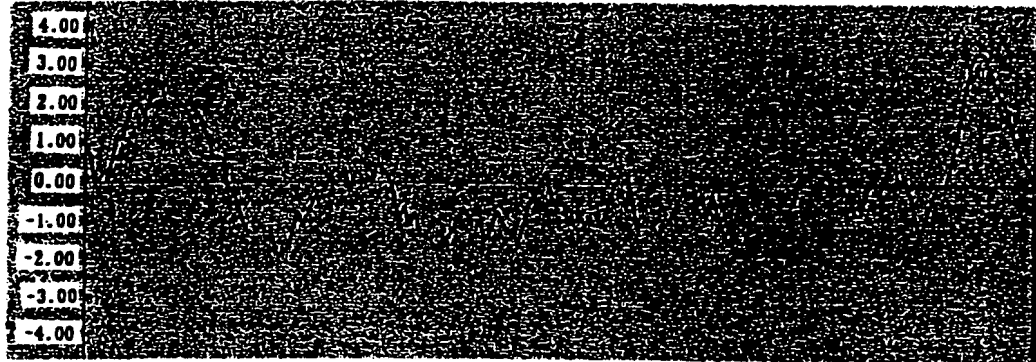


Fig. 5B

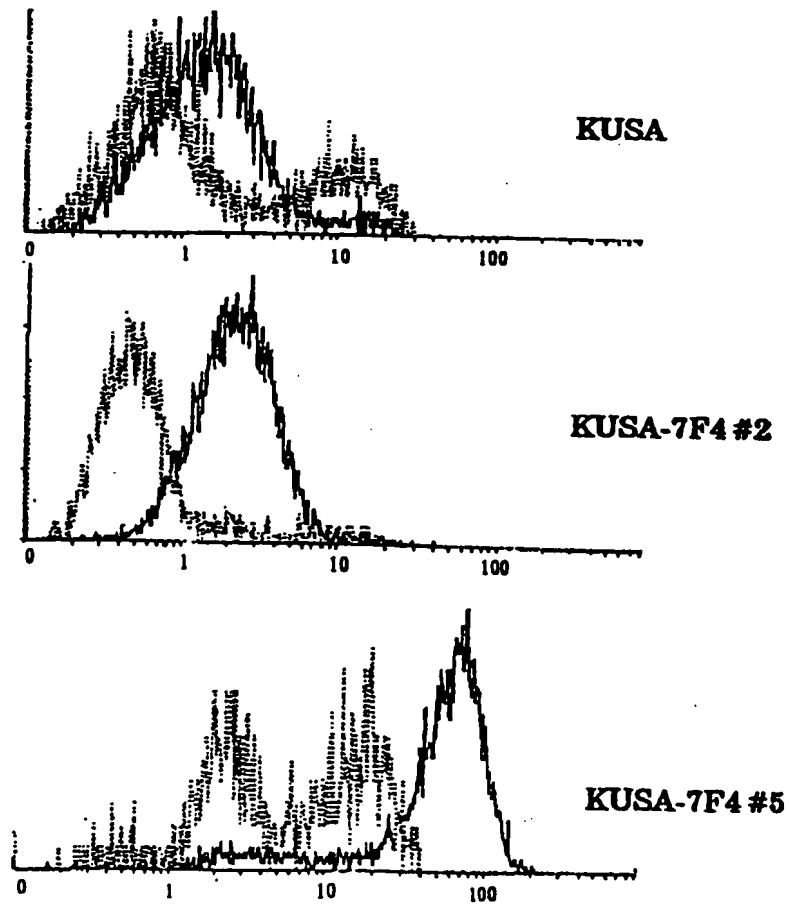


Fig. 6

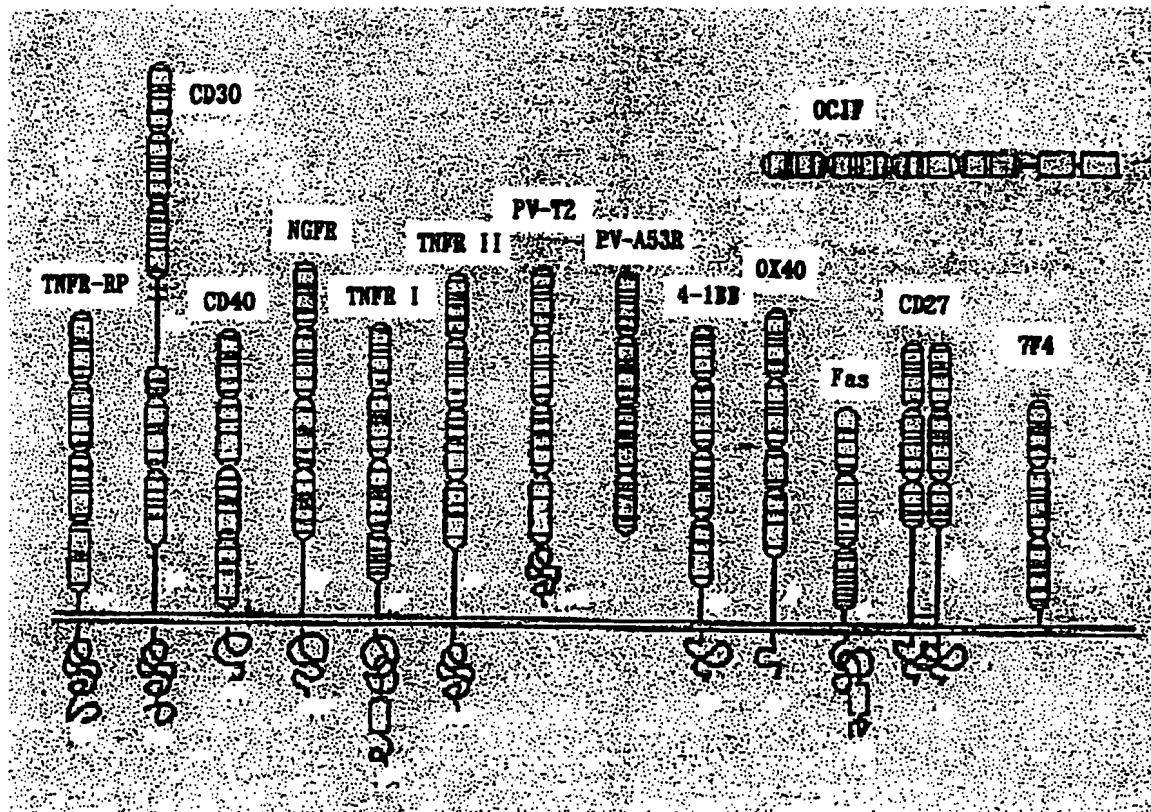


Fig. 7

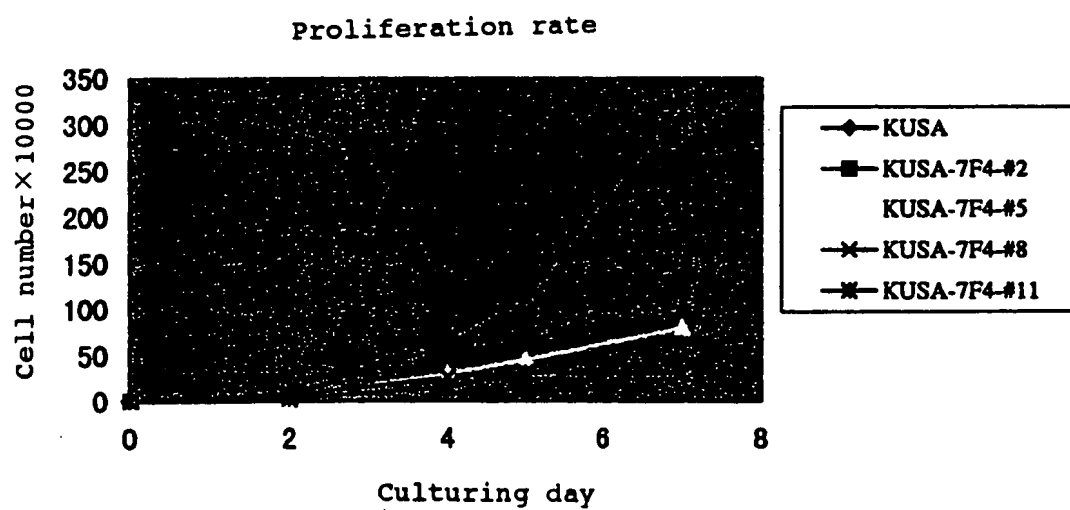


Fig. 8

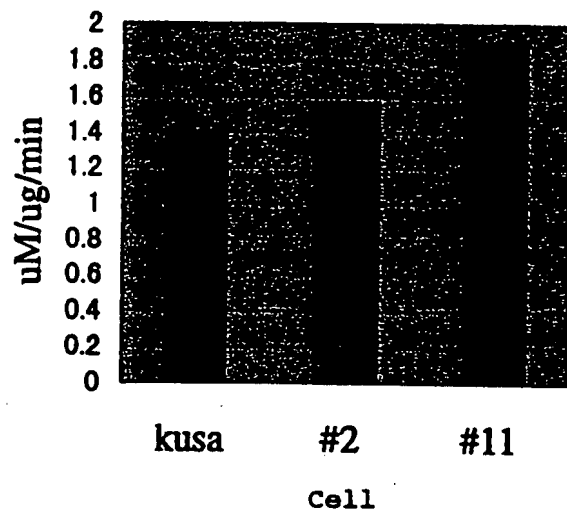




Fig. 9

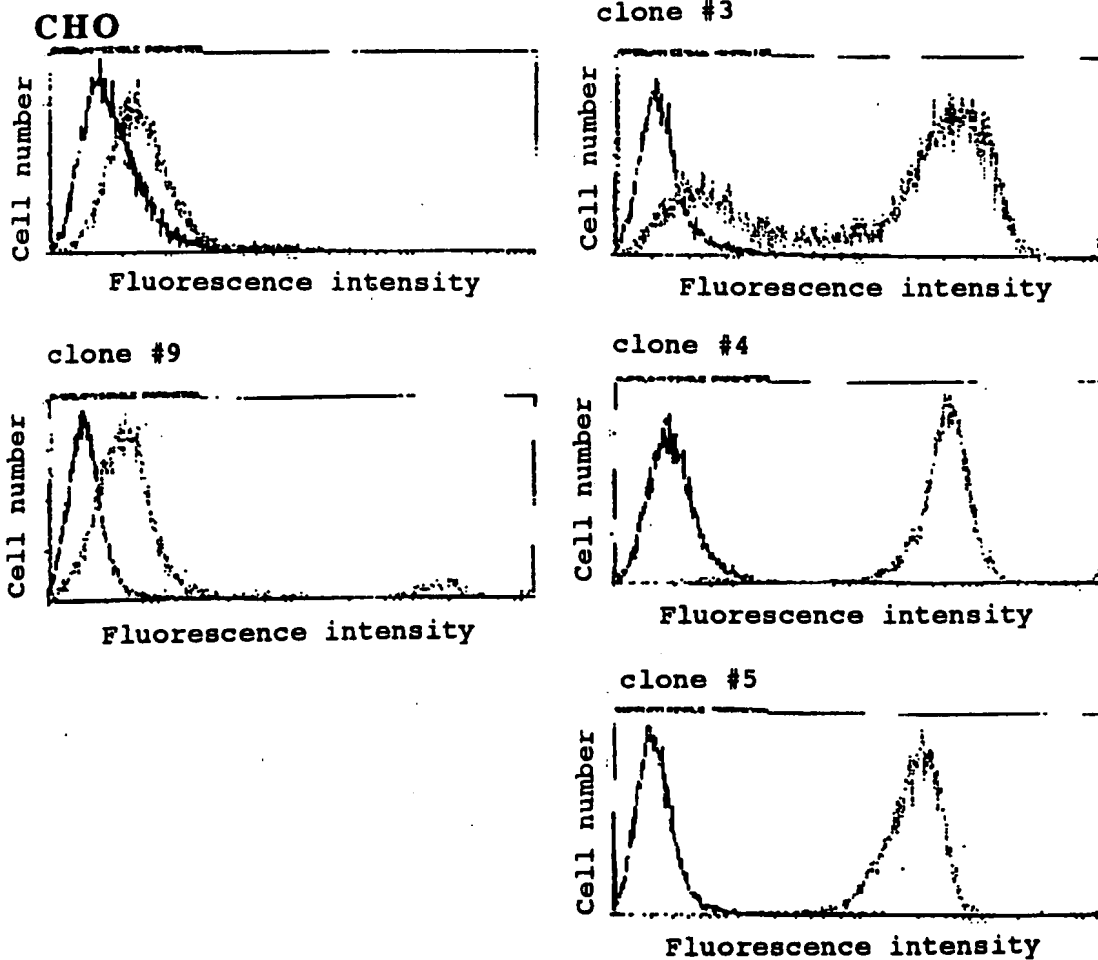


Fig. 10

